STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

P4r

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: *10/540,390*
Source: *P4r*
Date Processed by STIC: *2/24/06*

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
        http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , **EFS Submission User Manual** - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

| ERROR DETECTED | SUGGESTED CORRECTION | SERIAL NUMBER: _10/540,390_ |

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____Wrapped Nucleics Wrapped Aminos — The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2____Invalid Line Length — The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

3____Misaligned Amino Numbering — The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.

4____Non-ASCII — The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5____Variable Length — Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____PatentIn 2.0 "bug" — A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7____Skipped Sequences (OLD RULES) — Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8____Skipped Sequences (NEW RULES) — Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence.
<210> sequence id number
<400> sequence id number
000

9____Use of n's or Xaa's (NEW RULES) — Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.

10____Invalid <213> Response — Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence

11____Use of <220> — Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32)   (Sec. 1.823 of Sequence Rules)

12____PatentIn 2.0 "bug" — Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____ Misuse of n/Xaa — "n" can **only** represent a single <u>nucleotide</u>; "Xaa" can **only** represent a single <u>amino acid</u>

AMC – Biotechnology Systems Branch – 09/09/2003

PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/540,390

DATE: 02/24/2006
TIME: 12:56:10

Input Set : A:\21415-0019.txt
Output Set: N:\CRF4\02242006\J540390.raw

```
 3 <110> APPLICANT: Baell, Jonathan
 4        Huang, David
 5        Smith, Brian J
 6        Street, Ian P
 8 <120> TITLE OF INVENTION: Peptides and Therapeutic Uses Therefor
10 <130> FILE REFERENCE: 21415-0019
12 <140> CURRENT APPLICATION NUMBER: US 10/540,390
C--> 13 <141> CURRENT FILING DATE: 2005-06-23
15 <150> PRIOR APPLICATION NUMBER: AU 2002953561
16 <151> PRIOR FILING DATE: 2002-12-24
18 <150> PRIOR APPLICATION NUMBER: PCT/AU2003/001724
19 <151> PRIOR FILING DATE: 2003-12-24
21 <160> NUMBER OF SEQ ID NOS: 69
23 <170> SOFTWARE: PatentIn version 3.3
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 8
27 <212> TYPE: PRT
28 <213> ORGANISM: Artificial Sequence
30 <220> FEATURE:
31 <223> OTHER INFORMATION: Formula I sequence n=0, m=1
34 <220> FEATURE:
35 <221> NAME/KEY: MISC_FEATURE
36 <222> LOCATION: (1)..(1)
37 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
39 <220> FEATURE:
40 <221> NAME/KEY: MISC_FEATURE
41 <222> LOCATION: (2)..(2)
42 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
43        residue 3 or 7 by a linker
45 <220> FEATURE:
46 <221> NAME/KEY: MISC_FEATURE
47 <222> LOCATION: (3)..(3)
48 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
49        residue 2 or 7 by a linker
51 <220> FEATURE:
52 <221> NAME/KEY: MISC_FEATURE
53 <222> LOCATION: (4)..(4)
54 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
56 <220> FEATURE:
57 <221> NAME/KEY: MISC_FEATURE
58 <222> LOCATION: (5)..(5)
59 <223> OTHER INFORMATION: X = small amino acid residue
```

Does Not Comply
Corrected Diskette Needed

pp 1, 3-8

what is the source of genetic material?
(see item 11 on Error Summary Sheet)

61 <220> FEATURE:

```
 62 <221> NAME/KEY: MISC_FEATURE
 63 <222> LOCATION: (6)..(6)
 64 <223> OTHER INFORMATION: X = acidic amino acid residue
 66 <220> FEATURE:
 67 <221> NAME/KEY: MISC_FEATURE
 68 <222> LOCATION: (7)..(7)
 69 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
 70       residues 2 or 3 by a linker
 72 <220> FEATURE:
 73 <221> NAME/KEY: MISC_FEATURE
 74 <222> LOCATION: (8)..(8)
 75 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
 77 <400> SEQUENCE: 1
```

W--> 79 **Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa**

```
 80 1                5
 83 <210> SEQ ID NO: 2
 84 <211> LENGTH: 12
 85 <212> TYPE: PRT
 86 <213> ORGANISM: Artificial Sequence
 88 <220> FEATURE:
 89 <223> OTHER INFORMATION: Formula I sequence n=1, m=1
 92 <220> FEATURE:
 93 <221> NAME/KEY: MISC_FEATURE
 94 <222> LOCATION: (1)..(1)
 95 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
 97 <220> FEATURE:
 98 <221> NAME/KEY: MISC_FEATURE
 99 <222> LOCATION: (2)..(2)
100 <223> OTHER INFORMATION: X = small amino acid residue
102 <220> FEATURE:
103 <221> NAME/KEY: MISC_FEATURE
104 <222> LOCATION: (3)..(3)
105 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
106       residue 4, 6, 7 or 11 by a linker
108 <220> FEATURE:
109 <221> NAME/KEY: MISC_FEATURE
110 <222> LOCATION: (4)..(4)
111 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
112       residue 3, 6, 7 or 11 by a linker
114 <220> FEATURE:
115 <221> NAME/KEY: MISC_FEATURE
116 <222> LOCATION: (5)..(5)
117 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
119 <220> FEATURE:
120 <221> NAME/KEY: MISC_FEATURE
121 <222> LOCATION: (6)..(6)
122 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
123       residue 3, 4, 7 or 11 by a linker
125 <220> FEATURE:
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what is source? (handwritten annotation)

RAW SEQUENCE LISTING   DATE: 02/24/2006
PATENT APPLICATION: US/10/540,390   TIME: 12:56:10

Input Set : A:\21415-0019.txt
Output Set: N:\CRF4\02242006\J540390.raw

```
126 <221> NAME/KEY: MISC_FEATURE
127 <222> LOCATION: (7)..(7)
128 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
129       residue 3, 4, 6 or 11 by a linker
131 <220> FEATURE:
132 <221> NAME/KEY: MISC_FEATURE
133 <222> LOCATION: (8)..(8)
134 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
136 <220> FEATURE:
137 <221> NAME/KEY: MISC_FEATURE
138 <222> LOCATION: (9)..(9)
139 <223> OTHER INFORMATION: X = small amino acid residue
141 <220> FEATURE:
142 <221> NAME/KEY: MISC_FEATURE
143 <222> LOCATION: (10)..(10)
144 <223> OTHER INFORMATION: X = acidic amino acid residue
146 <220> FEATURE:
147 <221> NAME/KEY: MISC_FEATURE
148 <222> LOCATION: (11)..(11)
149 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
150       residue 3, 4, 6 or 7 by a linker
152 <220> FEATURE:
153 <221> NAME/KEY: MISC_FEATURE
154 <222> LOCATION: (12)..(12)
155 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
157 <400> SEQUENCE: 2
W--> 159 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
160 1               5                  10
163 <210> SEQ ID NO: 3
164 <211> LENGTH: 8
165 <212> TYPE: PRT
166 <213> ORGANISM: Artificial Sequence
168 <220> FEATURE:
169 <223> OTHER INFORMATION: Formula I sequence n=1, m=0
172 <220> FEATURE:
173 <221> NAME/KEY: MISC_FEATURE
174 <222> LOCATION: (1)..(1)
175 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
177 <220> FEATURE:
178 <221> NAME/KEY: MISC_FEATURE
179 <222> LOCATION: (2)..(2)
180 <223> OTHER INFORMATION: X = small amino acid residue
182 <220> FEATURE:
183 <221> NAME/KEY: MISC_FEATURE
184 <222> LOCATION: (3)..(3)
185 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
186       residue 4, 6, or 7 by a linker
188 <220> FEATURE:
189 <221> NAME/KEY: MISC_FEATURE
```

Input Set : **A:\21415-0019.txt**
Output Set: **N:\CRF4\02242006\J540390.raw**

```
     190 <222> LOCATION: (4)..(4)
     191 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
     192       residue 3, 6 or 7 by a linker
     194 <220> FEATURE:
     195 <221> NAME/KEY: MISC_FEATURE
     196 <222> LOCATION: (5)..(5)
     197 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
     199 <220> FEATURE:
     200 <221> NAME/KEY: MISC_FEATURE
     201 <222> LOCATION: (6)..(6)
     202 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
     203       residue 3, 4 or 7 by a linker
     205 <220> FEATURE:
     206 <221> NAME/KEY: MISC_FEATURE
     207 <222> LOCATION: (7)..(7)
     208 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is optionally linked
to
     209       residue 3, 4 or 6 by a linker
     211 <220> FEATURE:
     212 <221> NAME/KEY: MISC_FEATURE
     213 <222> LOCATION: (8)..(8)
     214 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
     216 <400> SEQUENCE: 3
W--> 218 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     219 1               5
     222 <210> SEQ ID NO: 4
     223 <211> LENGTH: 9
     224 <212> TYPE: PRT
     225 <213> ORGANISM: Artificial Sequence
     227 <220> FEATURE:
     228 <223> OTHER INFORMATION: Formula II sequence, m=0
     231 <220> FEATURE:
     232 <221> NAME/KEY: MISC_FEATURE
     233 <222> LOCATION: (1)..(1)
     234 <223> OTHER INFORMATION: X = any amino acid residue wherein it is linked to residue 8
by a
     235       linker
     237 <220> FEATURE:
     238 <221> NAME/KEY: MISC_FEATURE
     239 <222> LOCATION: (2)..(2)
     240 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
     242 <220> FEATURE:
     243 <221> NAME/KEY: MISC_FEATURE
     244 <222> LOCATION: (3)..(3)
     245 <223> OTHER INFORMATION: X = small amino acid residue
     247 <220> FEATURE:
     248 <221> NAME/KEY: MISC_FEATURE
     249 <222> LOCATION: (4)..(4)
     250 <223> OTHER INFORMATION: X = any amino acid residue
     252 <220> FEATURE:
     253 <221> NAME/KEY: MISC_FEATURE
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/540,390

DATE: 02/24/2006
TIME: 12:56:10

Input Set : A:\21415-0019.txt
Output Set: N:\CRF4\02242006\J540390.raw

```
254 <222> LOCATION: (5)..(5)
255 <223> OTHER INFORMATION: X = any amino acid residue
257 <220> FEATURE:
258 <221> NAME/KEY: MISC_FEATURE
259 <222> LOCATION: (6)..(6)
260 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
262 <220> FEATURE:
263 <221> NAME/KEY: MISC_FEATURE
264 <222> LOCATION: (7)..(7)
265 <223> OTHER INFORMATION: X = any amino acid residue
267 <220> FEATURE:
268 <221> NAME/KEY: MISC_FEATURE
269 <222> LOCATION: (8)..(8)
270 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is linked to residue
1 by
271        a linker
273 <220> FEATURE:
274 <221> NAME/KEY: MISC_FEATURE
275 <222> LOCATION: (9)..(9)
276 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
278 <400> SEQUENCE: 4
W--> 280 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
281 1                 5
284 <210> SEQ ID NO: 5
285 <211> LENGTH: 13
286 <212> TYPE: PRT
287 <213> ORGANISM: Artificial sequence
289 <220> FEATURE:
290 <223> OTHER INFORMATION: Formula II sequence, m=1
293 <220> FEATURE:
294 <221> NAME/KEY: MISC_FEATURE
295 <222> LOCATION: (1)..(1)
296 <223> OTHER INFORMATION: X = any amino acid residue, wherein it is linked to residue
8 by
297        a linker
299 <220> FEATURE:
300 <221> NAME/KEY: MISC_FEATURE
301 <222> LOCATION: (2)..(2)
302 <223> OTHER INFORMATION: X = hydrophobic amino acid residue
304 <220> FEATURE:
305 <221> NAME/KEY: MISC_FEATURE
306 <222> LOCATION: (3)..(3)
307 <223> OTHER INFORMATION: X = small amino acid residue
309 <220> FEATURE:
310 <221> NAME/KEY: MISC_FEATURE
311 <222> LOCATION: (4)..(4)
312 <223> OTHER INFORMATION: X = any amino acid residue
314 <220> FEATURE:
315 <221> NAME/KEY: MISC_FEATURE
316 <222> LOCATION: (5)..(5)
317 <223> OTHER INFORMATION: X = any amino acid residue
```

please correct this type of error in subsequent sequences

another example: "Peptide 17" in Sequence 17, as explanation for <213> Artificial Sequence

```
<210>  37                    Sequence
<211>  13
<212>  PRT
<213>  Artificial  Seqquence

<220>
<223>  Peptide 37    insufficient explanation - give source
<400>  37
```

Ile Ala Gln Glu Leu Arg Arg Ile Gly Asp Glu Phe Asn
1   5   10

```
<210>  38                    Sequence
<211>  26
<212>  PRT
<213>  Artificial  Sequuence

<220>
<223>  Peptide 38    insufficient
<400>  38
```

Asp Leu Arg Pro Glu Ile Arg Ile Ala Gln Glu Leu Arg Arg Ile Gly
1   5   10   15

Input Set : **A:\21415-0019.txt**
Output Set: **N:\CRF4\02242006\J540390.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8
Seq#:2; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12
Seq#:3; Xaa Pos. 1,2,3,4,5,6,7,8
Seq#:4; Xaa Pos. 1,2,3,4,5,6,7,8,9
Seq#:5; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13
Seq#:6; Xaa Pos. 1,2,3,4,5,6,7,8,9,10
Seq#:7; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14
Seq#:8; Xaa Pos. 1,2,3,4,5,6,7,8,9
Seq#:9; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12
Seq#:10; Xaa Pos. 1,2,3,4,5,6,7,8,9
Seq#:11; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13
Seq#:12; Xaa Pos. 1,2,3,4,5,6,7,8,9,10
Seq#:13; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14
Seq#:14; Xaa Pos. 1,2,3,4,5,6,7,8,9
Seq#:15; Xaa Pos. 1,8
Seq#:16; Xaa Pos. 4,11
Seq#:17; Xaa Pos. 4,11
Seq#:18; Xaa Pos. 4,11
Seq#:19; Xaa Pos. 5,12
Seq#:20; Xaa Pos. 4,11
Seq#:21; Xaa Pos. 4,11
Seq#:22; Xaa Pos. 4,11
Seq#:23; Xaa Pos. 4,11
Seq#:24; Xaa Pos. 2,4,11
Seq#:25; Xaa Pos. 4,11
Seq#:26; Xaa Pos. 4,11
Seq#:27; Xaa Pos. 4,11
Seq#:28; Xaa Pos. 4,11
Seq#:29; Xaa Pos. 4,11
Seq#:30; Xaa Pos. 4,11
Seq#:31; Xaa Pos. 4,11
Seq#:32; Xaa Pos. 4,11
Seq#:33; Xaa Pos. 1,8
Seq#:34; Xaa Pos. 4,11
Seq#:35; Xaa Pos. 4,11
Seq#:36; Xaa Pos. 4,11
```

VERIFICATION SUMMARY   DATE: 02/24/2006
PATENT APPLICATION: **US/10/540,390**   TIME: 12:56:11

Input Set : **A:\21415-0019.txt**
Output Set: **N:\CRF4\02242006\J540390.raw**

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:218 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:362 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:655 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:717 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:799 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:866 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:953 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:1019 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:1046 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:1073 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:1100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:1127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:1154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:1181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:1208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:1235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:1262 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:1294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:1321 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:1348 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:1375 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:1402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:1429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
L:1456 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:1483 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:1510 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:1537 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
L:1564 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
L:1591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:1618 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0
L:1625 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:37
L:1639 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:38